

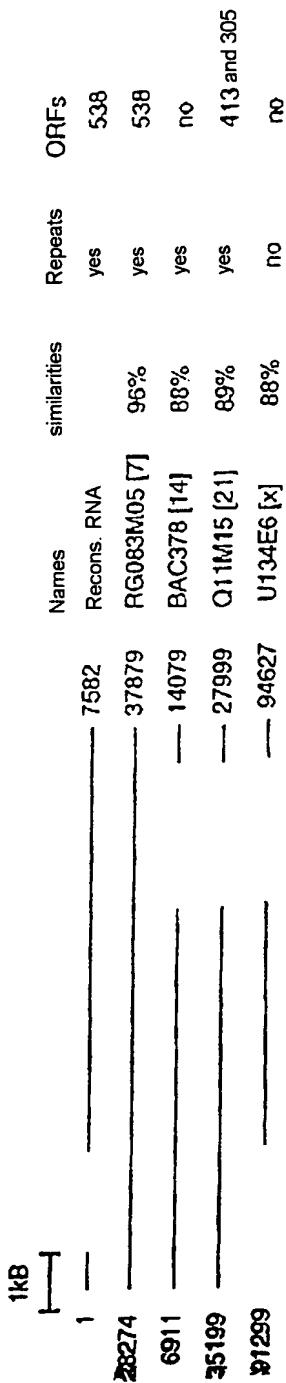
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WO 00/43521

PCT/FR00/00144

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FIG. 1



REPLACEMENT SHEET (RULE 26)

FIG 2

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	50
OCTAGAACGT ATTCTGGAGA ATTGGGACCA ATGTGACACT CAGACGCTAA					
P R T Y	S G E	L G P M	. H S	D A K	
L E R	I L E N	W D Q	C D T	Q T L R	
. N V	F W R	I G T N	V T L	R R	
GAAAGAAACG ATTTATATTTC TCTCTGAGT CCGCTGGCC ACAATATCCT					100
K E T	I Y I L	L Q Y	R L A	T I S S	
K K R	F I F	F C S T	A W P	Q Y P	
E R N D	L Y S	S A V	P P G H	N I L	
CTTCAAGGGA GAGAAACCTG CCTTCCTGAG GGAAGTATAAA ATTATAACAT					150
S R E	R N L A S	. G K Y K L	. H		
L Q G R	E T W L	P E G S I N	Y N I		
F K G	E K P G	F L R E V	. I I T S		
CATCTTACAG CTAGACCTCT TCTGTAGAAA GGAGGGAAA TGGAGTGAAG					200
H L T A	R P L L	. K G G Q M	E . S		
I L Q	L D L F	C R K E G K	W S E V		
S Y S	. T S S V E R	R A N G V K			
TGGCTATGTT GCAAACCTTC TTTTCAATTAA GAGAGAACTC ACAATTATGTT					250
A I C A N	F I K R Q L	T I M			
P Y V	Q T F	F S L R	D N S Q L C		
C H M C	K L S F H	. E T T H	N Y V		
AAAAAGTGIG GTTATGCCC TACAGGAAC CCTCAAGAGTC CACCTCCCTA					300
K V W	F M P	Y R K P	S E S T S L		
K K C G	L C P	T G S	P Q S P	P P Y	
K S V	V Y A L	Q E A	L R V	H L P T	
CCCCAGGTC CCTCTCCCCA CCTCTCTCTC AACTAAATAAG GACCCCCCTT					350
P Q R P	L P D	S F L N	. . G	P P F	
P S V	P S P T	P S S	T N K	D P P L	
P A S	P P R	L L P Q	L I R	T P L	
TAACCCAAAC GGCTCCAAAAG GAGATAGACA AAGGGTAAA CAATGAACCA					400
N P N	G P K G	D R Q	R G K	Q . T K	
T Q T	V Q K	E I D K	G V N	N E P	
. P K R	S K R R	R . T K G	. T	M N Q	
AAGAGTGOCA ATATTCGGG ATATGCCCC CCTCAAGAG TGAGAGGAGG					450
E C Q	Y S P	I M P P	P S S	E R R	
K S A N	I P R	L C P	L Q A V	R G G	
R V P	I F P D	Y A P	S K Q	. E E E	
AGAATTCGGC CCAGCCAGAG TGGCTGTACCT TTTTCTCTTC TCAGACTTAA					500
R I R P	S Q S	A C T	F F S L	R L K	
E F G	P A R V	P V P	F S L	S D L K	
N S A	Q P E	C L Y L	F L S	Q T .	

REPLACEMENT SHEET (RULE 26)

## FIG 2 (continued)

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	1234567890
ACCAAAATTAA	AATAGACCTA	CGTAAATTCT	CAGATAACCC	TGACGGCTAT	550
A N .	N R P R .	I L R .	P .	R L Y	
Q I K I	I D L G K F S	D N P	D G Y		
S K L K .	T .	V N S Q I T L	T A I		
ATTGATGTTT	TACAAGGGTT	AGGACAATCC	TTTGTACTGA	CATGGAGAGA	600
. C F	T R V R T I L	. S D M E R			
I D V L	Q G L G Q S	F D L T	W R D		
L M F Y K G .	D N P L I .	H G E I			
TATAATGTTA	CTACTAAATTC	AGACACTAAC	CCCAAATGAG	AGAAGTGGCG	650
Y N V T	T K S D T N P K .	E K C R			
I M L L L N Q	T L T P N E R S A A				
. C Y Y .	I R H .	P Q M R E V P			
CTGTAACCTGC	AGCCCCGAGAG	TTTCCCCGATC	TTTGGTATCT	CAGTCAGGCC	700
C N C S P R V	W R S L V S	Q S G Q			
V T A A R E	F G D L W Y L S Q A				
L . L Q P E S	L A I F G I S . V R P				
AACAATAGGA	TGACAACAGA	GGAAAAGAAC	ACTCCCCACAG	GCCAGCCAGGC	750
Q . D D N R	G K N N S H R	P A G			
N N R M T T E	E R T T P T G	Q Q A			
T I G . Q Q R	K E Q L P Q	A S R Q			
AGTTCCCCAGT	GTAGACCCCT	ATTGGGACAC	AGAATAGAA	CATGGAGATT	800
S S Q C R P S	L G H R I R T	W R L			
V P S V D P H	W D T E S E	H G D W			
F P V . T L I	G T Q N Q N	M E I			
GGGCCACAA	ACATTTGCCTA	ACTTGCGTC	TAGAAGGACT	GAGGAAACT	850
V P Q T F A N	L R A R R T	E E N .			
C H K H L L T	C V L E G L R K T				
G A T N I C .	L A C . K D .	G K L			
AGGAAGAAC	CTATGAATTA	CTCAATGATG	TOCACATATAA	CACAGGGAAA	900
E E A Y E L	L N D V H Y N T	G K			
R K K P M N Y	S M M S T I T	Q G K			
G R S L . I T	Q . C P L .	H R E R			
CGAAGAAAAT	CTTACTGCTT	TCTGGACAG	ACTAAGGGAG	GCATTGAGGA	950
G R K S Y C F	S G Q T K G G	I E E			
E E N L T A F	L D R L R E	A L R K			
K K I L L L F	W T D . G R H .	G			
AGCATACCTC	CCGTGTCACCT	GACTCTATTC	AAAGGCCAACT	AACTCTAAAG	1000
A Y L P V T .	L Y .	R P T N L K G			
H T S L S P	D S I E	G Q L I L K			
S I P P C H L	T L L K A N .	S . R			

FIG 2 (continued)

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	1050
GATAAGTTTA TCACTCAGTC AGCTGCAGAC ATTAGAAAAAA ACTTCAAAAG					
. V Y H S V S C R H . K K L Q K					
D K F I T Q S A A D I R K N F K S					
I S L S L S Q L Q T L E K T S K V					
TCTGCCTTAC GCGCGGACCA GAACTTAGAA ACCGTTATTA ACTTGGCAGC					1100
S A L G P E Q N L E T L F N I A S					
L P . A R S R T . K P Y L T W H P					
C L R P G A E L R N P I . L G I					
CTCAGTTTT TATAATAGAG ATCAGGAGGA GCACCGAAA CGGGACAAAC					1150
S V F Y N R D Q E E Q A K R D K R					
Q F F I I E I R R S R R N G T N					
L S F L . . R S G G A G E T G Q T					
CGGATAAAA AAAAAGGGG GGTCCACTAC TTTAGTCATG CCCTCAAGC					1200
D K K K R G G P L L . S W P S G					
G I K K K G G V H Y F S H G P Q A					
G . K K K G G S T T L V M A L R Q					
AAGCAGACTT TGGACCCCTCT GCAAAAGGG AAGGCTGGCC AATTCAAATG					1250
K Q T L E A L Q K G K A G Q I K C					
S R L W R L C K R E K L G K S N A					
A D F G G S A K G K S W A N Q M					
CTTAATAGG CTCGCTTCA GCGCGCTCA CAAGGACACT TTAAAAAAAAGA					1300
L I G L A S S A V Y K D T L K K I					
. . G W L P V R S T R T L . K R					
P N R A G F Q C G L Q G H F K K D					
TTCATCCAAGT AGAAATAAGC CGCGCGCTTG TOCATCGCC TTACGTCAAG					1350
I Q V E I S R P L V H A P Y V K					
L S K . K . A A P L S M P L T S R					
Y P S R N K P P P C P C P L R Q G					
GGAATCACTG GAAGGCGCAC TGCGCGAGG GATGAAGATA CTCTGAGTCA					1400
G I T G R P T A P G D E D T L S Q					
E S L E G P L P Q G M K I L . V R					
N H W K A H C P R G . R Y S E S					
GAAGCCATTA ACCAGATGAT CGAGCAGAG GACTGAGGGT CGGGGGGGCG					1450
K P L T R . S S S R T E G A R G E					
S H . P D D P A A G L R V P G A					
E A I N Q M I Q Q Q D . G C P G R					
AGGGCGAGCC CATGCCATCA CGCTCAGAGA CGGGGGGGTA TGTTTGACCA					1500
R Q P M P S P S Q S P G Y V . P					
S A S P C H H P H R A P G M F D H					
A P A H A I T L T E P R V C L T I					

## FIG 2 (continued)

10	20	30	40	50
1234567890	1234567890	1234567890	1234567890	1234567890
<u>TTGAGACCCA A</u>				
L	R	A		1511
.	E	P		
	E	S	Q	